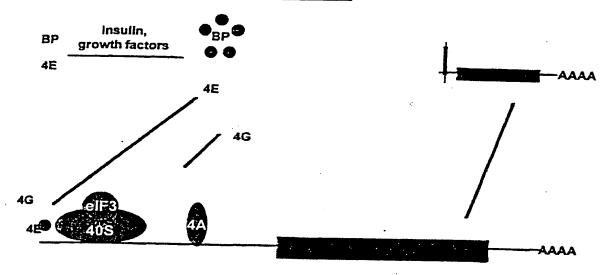


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### FIGURE 4



h4g human eiF4G BβKKRYDREFLLGFAARQIKIWFQNRRMKWKK
S scrambled eiF4G BβFDLKFALGRYRAEKRQIKIWFQNRRMKWKK
- no peptide

rec 4E recombinant human elF4E

Peptide_	-		W	1	2	
elF4E-		****				

H human elF4G B β β K K R Y D R E F L L G F 413-424
Y yeast elF4G B β β K Y T Y G P T F L L Q F 449-460
W wheat elF4G B β β R V R Y S R D Q L L D L 62-73
1 human 4E-BP1 B β β R I I Y D R K F L M E U 51-62
2 human 4E-BP2 B β β R I I Y D R K F L L D R 51-62
S scrambled elF4G

#### FIGURE 6

S H<sub>wt</sub> H<sub>3A</sub> W<sub>wt</sub> W<sub>3A</sub> BP1 - + - + - + - +

-elF4E

4G Peptide	Sequence													
H <sub>wt</sub> hu 4G <sub>(413-424)</sub>	K	K	R	Y	D	R	E	F	L	L	G	F	A	A
H <sub>3A</sub> hu 4G <sub>(413-424)YALALA</sub>	K	K	R	A	D	R	E	F	A	A	G	F	A	A
W <sub>wt</sub> wh 4G <sub>(62-73)</sub>	R	V	R	Y	5	R	D	Q	L	L	D	L	A	A
W3A wh 4G(82-73)YALALA														A
S scrambled hu 4G	F	D	L	K	F	A	L	G	R	Y	R	A	E	K

all peptides biotinylated and linked to Penetratin™

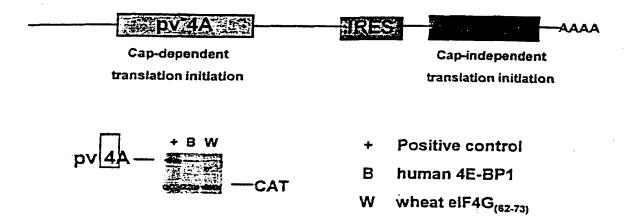
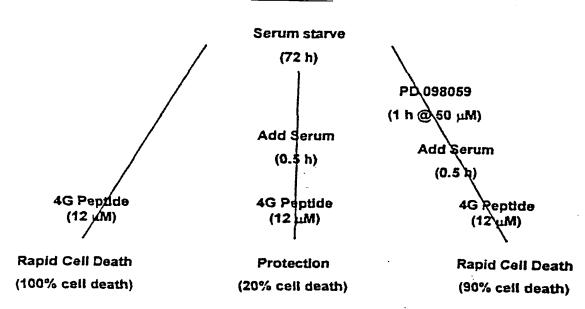


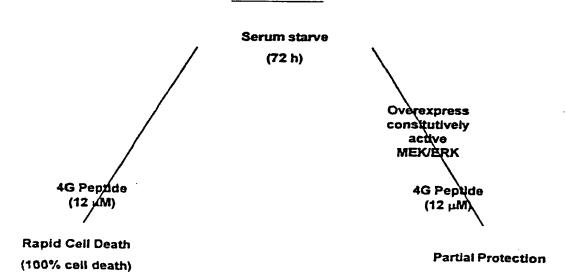
FIGURE 8

Peptide	Concentration							
replide	ЗμМ	6 μ <b>M</b>	9 μΜ	12 µM				
hu 4G <sub>(413-424)</sub>	+	++	+++	+++				
hu 4G <sub>(413-424)Y416AL421AL422A</sub>	-	•		-				
wh 4G <sub>(62-73)</sub>	-/+	+	++	+++				
wh 4G(62-73)Y65AL70AL71A	-	•	-	<b>-</b>				
scrambled hu 4G	_		•	-				

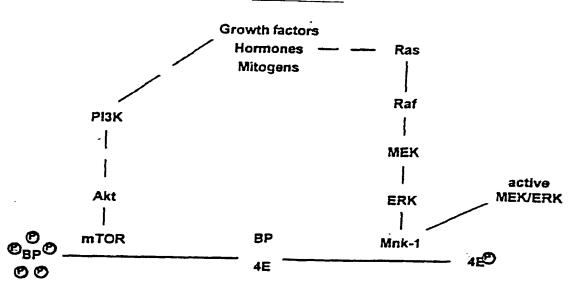
MRC5 cells, 72 h serum-free growth, all peptides blotinylated and linked to PenetratinTM





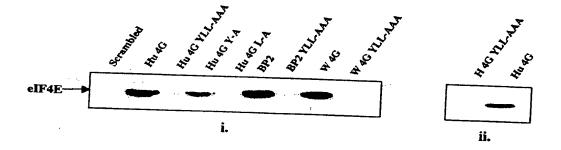




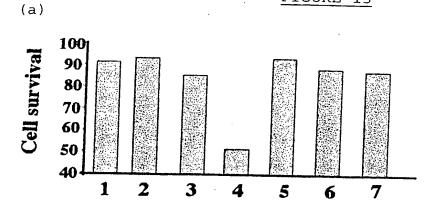


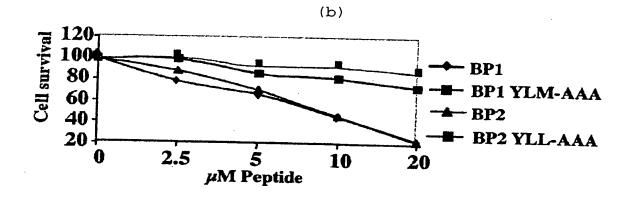
(a)			
Hu	4G 4G YLL-AAA 4G Y-A 4G L-A	Human eIF4G Peptide (569-580)Wild Type Human eIF4G Peptide (569-580)Y572A L577A L578A Human eIF4G Peptide (569-580)Y572A Human eIF4G Peptide (569-580)L577A	KKRYDREFLLGF KKRADREFAAGF KKRADREFLLGF KKRYDREFALGF
	YLL-AAA	Wheat eIF4G Peptide (62-73)Wild Type Wheat eIF4G Peptide (62-73)Y65A, L70A, L71A	RVRYSRDQLLDL RVRASRDQAADL
	YLL-AAA	Human 4E-BP2 Peptide (51-62)Wild Type Human 4E-BP2 Peptide (51-62)Y54A, L59A, L60A.	RIIYDRKFLLDR RIIADRKFAADR
BP1 BP1	YLM-AAA	Human 4E-BP1 Peptide (51-62)Wild Type Human 4E-BP1 Peptide (51-62)Y54A, L59A, M60A	RIIYDRKFLMEU RIIADRKFAAEU

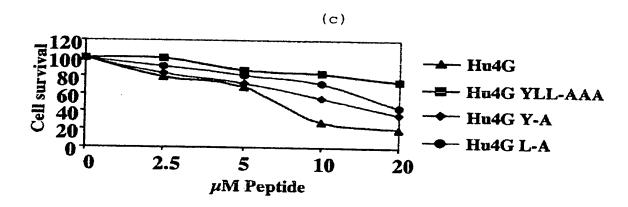
(b)

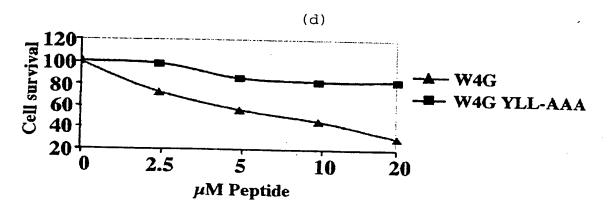




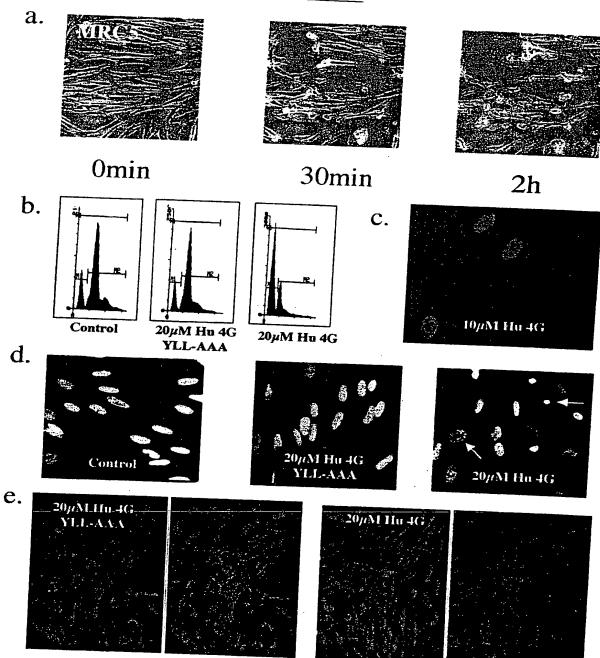


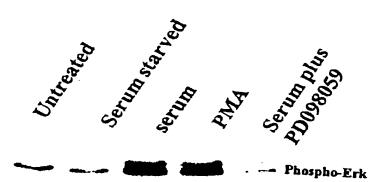




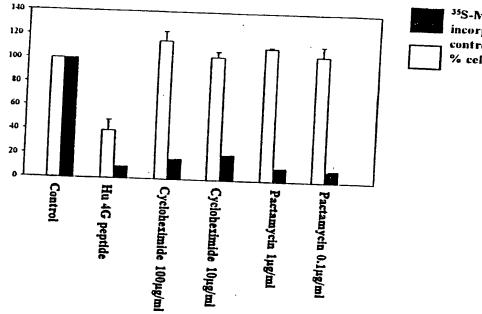


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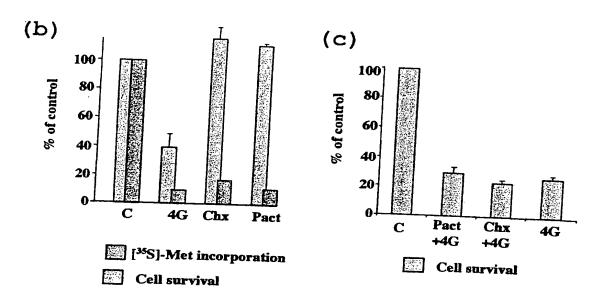


(a)



35S-Met
incorporation-% of
control
% cell survivat

## FIGURE 16(cont.)



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